## **AMENDMENTS**

## Amendments to the Claims:

This listing of claims will replace all prior versions, and listings, of claims in the application.

- 1. (**Currently amended**) An isolated FB005, FB006 or FB066 peptide comprising the sequence of SEQ ID NO:1, SEQ ID NO:2 or SEQ ID NO:3, respectively.
- 2. (Canceled)
- 3. (Canceled)
- 4. (Previously presented) An isolated, modified peptide selected from the group consisting of:
  - (a) SEQ ID NO:1;
  - (b) SEQ ID NO:2;
  - (c) SEQ ID NO:3; and
  - (d) SEQ ID NO:7,

and having at least one substituted amino acid residue at a predetermined position in the peptide sequence, wherein the at least one substituted amino acid residue is a hydrophilic amino acid residue, a hydrophobic amino acid residue, an amino acid residue having a propensity to form alpha helices, a D-isomer of one of the naturally occurring L-amino acids, or a non-naturally occurring amino acid residue.

- 5. (Previously presented) An isolated, derivatized peptide selected from the group consisting of:
  - (a) the FB005M peptide of SEQ ID NO:8;
  - (b) the FB005CM peptide of SEQ ID NO:9;
  - (c) the FB006M peptide of SEQ ID NO:10;
  - (d) the FB007M peptide of SEQ ID NO:11;
  - (e) the FB010M peptide of SEQ ID NO:12;

- (f) the FB010KM peptide of SEQ ID NO:13;
- (g) the FB066M peptide of SEQ ID NO:14; and
- (h) the FB066KM peptide of SEQ ID NO:15.
- 6. (Previously presented) An isolated, derivatized peptide selected from the group consisting of:
  - (a) SEQ ID NO:1;
  - (b) SEQ ID NO:2;
  - (c) SEQ ID NO:3; and
  - (d) SEQ ID NO:7,

wherein predetermined amino acid residues in the peptide sequence are derivatized by conjugating a coupling group to said predetermined amino acid residues.

- 7. (Previously presented) The modified peptide of claim 4, wherein predetermined amino acid residues in the peptide sequence are derivatized by conjugating a coupling group to said predetermined amino acid residues.
- 8. (Currently amended) The isolated peptide of claim <u>41</u>, wherein the peptide <u>is SEQ</u> <u>ID NO:1 and</u> is derivatized by attaching a coupling group to a lysine, said lysine being substituted for glutamic acid at position 23 or added at the C-terminus.
- 9. (Currently amended) The isolated peptide of claim <u>42</u>, wherein the peptide <u>is SEQ</u> <u>ID NO:2 and</u> is derivatized by attaching a coupling group to the lysine at position 13.
- 10. (Currently amended) The isolated peptide of claim 42, wherein the peptide is SEQ ID NO:2 and is modified by substituting the lysine at position 13 with glutamic acid and derivatized by attaching a coupling group to an additional lysine residue added at the C-terminus.
- 11. (Previously presented) An isolated, derivatized peptide consisting of the sequence of SEQ ID NO: 3, wherein the peptide is modified by replacing glutamic acid at position 13 with a lysine and attaching a coupling group to the lysine, or derivatized by conjugating a coupling group to a lysine added at the C-terminus.

- 12. (**Currently amended**) The isolated peptide of claim <u>43</u>, wherein the peptide <u>is SEQ</u> <u>ID NO:7 and</u> is derivatized by attaching a coupling group to the lysine at position 13, or to an additional lysine added at the C-terminus.
- 13. (Currently amended) The derivatized peptide of any one of claims 5 12 7, wherein the coupling group is selected from the group consisting of:
  - (a) a maleimido group;
  - (b) a succinimidyl group;
  - (c) a hydrazine group; and
  - (d) a carbonyl group.
- 14. (Previously presented) The derivatized peptide of claim 13, wherein the maleimido group is 3'-maleimidopropionate connected to the epsilon amino group of lysine by [2-(2-amino)ethoxyl]ethoxy acetic acid.
- 15. (Currently amended) A pharmaceutical composition comprising the peptide of any one of claims 1 or -4 or the derivatized peptide of any one of claims 5 12 7.
- 16. (**Currently amended**) A conjugate comprising the derivatized peptide of <del>any one of claims 5-12</del> conjugated to a blood component.
- 17. (Previously presented) The conjugate of claim 16, wherein the blood component is selected from the group consisting of:
  - (a) human serum albumin protein;
  - (b) human transferrin protein;
  - (c) human ferritin protein;
  - (d) human immunoglobulin proteins;
  - (e) human ferritin protein;
  - (f) human α-2-macroglobulin protein;
  - (g) human thyroxin binding protein;
  - (h) human steroid binding proteins; and
  - (i) combinations thereof.

- 18. (Currently amended) A method for <u>preventing or reducing infection of</u>, or <u>preventing viral replication in</u>, mammalian cells by a virus comprising presenting a peptide according to <u>any one of claims 1 or -4</u> or a peptide derivative according to <u>any one of claims 5 12 7</u> to said mammalian cells.
- 19. (Canceled)
- 20. (Canceled)
- 21. (Currently amended) The method of claims 18-20, wherein said peptide is presented in the presence of said virus.
- 22. (Currently amended) The method of claims 18-21, wherein the virus is selected from the group consisting of:
  - (a) human immunodeficiency virus (HIV); and
  - (b) simian immunodeficiency virus (SIV).
- 23. (**Currently amended**) The method of claims 18–21, wherein the peptide or peptide derivative is administered orally, topically, intravascularly, intravarterially, intramuscularly, or subcutaneously.
- 24. (Currently amended) The method of claims 18–21, wherein the peptide or peptide derivative is co-administered with one or more additional HIV treatment(s).
- 25. (Previously presented) The method of claim 24, wherein the said one or more additional HIV treatment(s) comprises at least one other variant gp41 peptide.
- 26. (Previously presented) The method of claim 24, wherein the additional HIV treatment(s) is selected from the group consisting of:
  - (a) AGENERASE;
  - (b) COMBIVIR;
  - (c) CRIXIVAN;
  - (d) EMTRIVA;
  - (e) EPIVIR;
  - (f) FORTOVASE;
  - (g) HIVID;

(h) INVIRAS	SE;
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- (i) KALETRA;
- (j) NORVIR;
- (k) RESCRIPTOR;
- (l) RETROVIR;
- (m) REYATAZ;
- (n) SUSTIVA;
- (o) TRIZIVIR;
- (p) VIDEX EC;
- (q) VIDEX;
- (r) VIRACEPT;
- (s) VIRAMUNE;
- (t) VIREAD;
- (u) ZERIT; and
- (v) ZIAGEN.
- 27. (**Currently amended**) The method of claims 18–21, wherein the virus is HIV and the mammalian cells are human cells.
- 28. (Currently amended) A method of preventing or reducing HIV infection comprising administering a derivative variant gp41 peptide of any one of claims 5-12 7 to a patient whose cells have been exposed to HIV, wherein said peptide derivative conjugates with a blood component of said patient, thereby extending the half-life of the peptide in said patient's blood.
- 29. (Previously presented) A method of making an antiviral conjugate comprising mixing derivatized variant gp41 peptide(s) with blood components and allowing the formation of covalent bonds between derivatized variant gp41 peptide and blood components.
- 30. (Currently amended) The method of claims 27–28, wherein the blood component is selected from the group consisting of:

- (a) human serum albumin protein;
- (b) human transferrin protein;
- (c) human ferritin protein;
- (d) human immunoglobulin proteins;
- (e) human ferritin protein;
- (f) human  $\alpha$ -2-macroglobulin protein;
- (g) human thyroxin binding protein;
- (h) human steroid binding proteins; and
- (i) combinations thereof.
- 31. (Previously presented) The method of claim 29, wherein the blood component is human serum albumin protein.
- 32. (**Currently amended**) The method of claims 27–28, wherein the conjugation occurs in vivo.
- 33. (Currently amended) The method of claims 27-28, wherein the conjugation occurs ex vivo.
- 34. (Previously presented) The method of claim 32, wherein the blood component(s) are separated by plasmaphoresis before conjugation to the derivatized peptide.
- 35. (**Currently amended**) A pharmaceutical composition comprising the isolated peptide of claims 1 or 4 14, or the derivatized peptide of claim 7, and a pharmaceutically acceptable carrier.
- 36. (Previously presented) A method for the generation of peptides having anti-viral, virostatic or anti-fusogenic activity comprising:
  - (a) screening a viral virulence protein(s) to identify sequences thereof having alphahelical forming propensities;
  - (b) designing an altered peptide by modifying or derivatizing at least one amino acid residue(s) of said identified sequence;
  - (c) synthesizing said altered peptides; and
  - (d) testing said peptides to verify anti-viral, virostatic or anti-fusogenic activity.